

Johnson

1806

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RAW SEQUENCE LISTING
PATENT APPLICATION US/08/436,339

DATE: 10/15/97
TIME: 12:39:25

INPUT SET: S21003.raw

#11
B. Denny
10/15/97

This Raw Listing contains the General
Information Section and up to the first 5 pages.

ENTERED

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Papayannopoulou, Thalia

(ii) TITLE OF INVENTION: Peripheralization of Hematopoietic
Stem Cells

(iii) NUMBER OF SEQUENCES: 2

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Allegretti & Witcoff, Ltd.
(B) STREET: 75 State Street
(C) CITY: Boston
(D) STATE: Massachusetts
(E) COUNTRY: USA
(F) ZIP: 02109

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 08/436,339
(B) FILING DATE: 13-JUL-1995
(C) CLASSIFICATION: 424

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 07/977,702
(B) FILING DATE: 13-NOV-1992

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Keown, Wayne A
(B) REGISTRATION NUMBER: 33,923
(C) REFERENCE/DOCKET NUMBER: 92,678; D014

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 617/345-9100
(B) TELEFAX: 617/345-9111

(2) INFORMATION FOR SEQ ID NO:1:

Replaced
By
Paper
#19
7/02/98

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/436,339DATE: 10/15/97
TIME: 12:39:28

INPUT SET: S21003.raw

47 (i) SEQUENCE CHARACTERISTICS:
48 (A) LENGTH: 360 base pairs
49 (B) TYPE: nucleic acid
50 (C) STRANDEDNESS: single
51 (D) TOPOLOGY: linear
52

53 (ii) MOLECULE TYPE: cDNA
54

55 (iii) HYPOTHETICAL: NO
56

57 (iv) ANTI-SENSE: NO
58

59 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
60

61
62 GTCAAAGTGC AGCAGTCTGG GGCAGAGCTT GTGAAGCCAG GGGCCTCAGT CAAGTTGTCC 60
63
64 TGCACAGCTT CTGGCTTCAA CATTAAAGAC ACCTATATGC ACTGGGTGAA GCAGAGGCCT 120
65
66 GAACAGGGCC TGGAGTGGAT TGAAGGATT GATCCTGCGA GTGGCGATAC TAAATATGAC 180
67
68 CCGAAGTTCC AGGTCAAGGC CACTATTACA GCGGACACGT CCTCCAACAC AGCCTGGCTG 240
69
70 CAGCTCAGCA GCCTGACATC TGAGGACACT GCCGTCTACT ACTGTGCAGA CGGAATGTGG 300
71
72 GTATCAACGG GATATGCTCT GGACTTCTGG GGCCAAGGGA CCACGGTCAC CGTCTCCTCA 360
73
74
75
76

77 (2) INFORMATION FOR SEQ ID NO:2:
78

79 (i) SEQUENCE CHARACTERISTICS:
80 (A) LENGTH: 318 base pairs
81 (B) TYPE: nucleic acid
82 (C) STRANDEDNESS: single
83 (D) TOPOLOGY: linear
84

85 (ii) MOLECULE TYPE: cDNA
86

87 (iii) HYPOTHETICAL: NO
88

89 (iv) ANTI-SENSE: NO
90

91 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
92

93
94 AGTATTGTGA TGACCOAGAC TCCCAAATTC CTGCTTGTTT GAGCAGGAGA CAGGGTTACC 60
95
96 ATAACTGCA AGGCCAGTCA GAGTGTGACT AATGATGTAG CTGGGTACCA ACAGAAGCCA 120
97
98 GGGCAGTCTC CTAAACTGCT GATATATTAT GCATCCAATC GCTACACTGG AGTCCCTGAT 180
99

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/436,339DATE: 10/15/97
TIME: 12:39:31

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102	GAAGACCTGG CAGTTTATTT CTGTCAGCAG GATTATAGCT CTCCGTACAC GTTCGGAGGG	300
103		
104	GGGACCAAGC TGGAGATC	318
105		
106		

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/436,339

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